Kaveri et al. 10/031,938

characterized in that it comprises a protease inhibitor. Examples of protease inhibitors that can be used as anti-Factor VIII allo-antibody catalysed Factor VIII degradation inhibitors within the context of the present invention, without being limited thereto, are fluorophosphate-type inhibitors, such as DFP for example, or sulphonyl fluoride-type inhibitors, such as PMSF or AEBSF (4-(2-aminoethyl)benzenesulphonyl fluoride hydrochloride (notably marked by Roche Diagnostics GmbH, Mannheim, Germany, under the trademark Pefabloc®)), for example. More particularly, this inhibitor is characterized in that said inhibitor inhibits cleavage of the scissile bonds: Arg<sup>372</sup>-Ser<sup>373</sup>, located between the A1 and A2 domains, Tyr<sup>1680</sup>-Asp<sup>1681</sup>, located on the N-terminus of the A3 domain, and the Glu<sup>1794</sup> – Asp<sup>1795</sup> located within the A3 domain of the Factor VIII molecule. More preferably still, this inhibitor is characterized in that it comprises a peptide or non-peptide analogue of the amino acid sequence:

Ser Val Ala Lys Lys His Pro;

a peptide or non-peptide analogue of the amino acid sequence:

Asp Glu Asp Glu Asn Gln Ser; or

a peptide or non-peptide analogue of the amino acid sequence:

Asp Gln Arg Gln Gly Ala Glu.

On page 20, please delete the existing table and replace it with the following table:

Amino acid sequence	Cleavage site
Ser Val Ala Lys Lys His Pro (SVAKKHP)  Asp Gln Arg Gln Gly Ala Glu (DQRQGAE)	$Arg_{372}^{372} - Ser_{373}^{373}$ $(R^{372} - S^{373})$ $Glu_{1794}^{1794} - Asp_{1795}^{1795}$ $(E^{1794} - D^{1795})$ $Tyr_{1680}^{1680} - Asp_{1681}^{1681}$
Asp Glu Asp Glu Asn Gln Ser (DEDENQS)	$(Y^{1680} - D^{1681})$

Doc. No.: 1154664 01/27/03 1:16 PM